- 1/1 (C) WPI / DERWENT
- AN 2001-610074 [70]
- AP JP20000060548 20000306
- PR JP20000060548 20000306
- TI New G protein-coupled receptor polypeptide for use in the development of new drugs
- IW NEW PROTEIN COUPLE RECEPTOR POLYPEPTIDE DEVELOP NEW DRUG
- PA (KYOW) KYOWA HAKKO KOGYO KK
- PN JP2001245666A-20010911 DW200170 C12N15/09 126pp
- IC A01H5/00; A01K67/027; A61K39/395; A61K45/00; A61P25/00; A61P35/00; C07K14/705; C07K16/28; C12N1/15; C12N1/19; C12N1/21; C12N5/10; C12N15/09; C12P21/02; C12P21/08; C12Q1/68; G01N33/15; G01N33/50; G01N33/53; G01N33/566
- AB JP2001245666 NOVELTY A G protein-coupled receptor (GPCR) polypeptide having a sequence (S1) of 371 amino acids, given in the specification, is new.
  - DETAILED DESCRIPTION INDEPENDENT CLAIMS are also included for the following:
  - (1) a polypeptide having an amino acid sequence in which at least one amino acid is deleted, replaced or added in S1 and having an activity substantially same as the above polypeptide;
  - (2) a partial peptide of the above GPCR polypeptide having combinability to a ligand, an agonist, an antagonist or a function-modifying substance of the polypeptide;
  - (3) a DNA encoding the above GPCR polypeptide;
  - (4) a DNA having a sequence of bases 175 to 1287 in a sequence of 1714 base pairs (bp), given in the specification;
  - (5) a DNA hybridizing with the above DNA under a stringent condition and encoding a polypeptide having an activity substantially the same as the above GPCR polypeptide;
  - (6) a DNA encoding a partial peptide of the above GPCR polypeptide;
     and
  - (7) a recombinant DNA prepared by recombining the above DNA to a vector; and
  - (8) a transformed cell, a transformed plant or a transformed nonhuman animal carrying the above recombinant DNA.
  - USE The GPCR polypeptide can be used for the development of new drugs.
  - (Dwg.0/16)

```
1: BD017045. Novel polypeptide...[gi:22558221]
```

## 10/539565 JC17 Rec'd PCT/PTO 17 JUN 2005 PAT 27-AUG-2002

LOCUS BD017045 1714 bp Novel polypeptide. DEFINITION BD017045 ACCESSION BD017045.1 GI:22558221 VERSION JP 2001245666-A/1. KEYWORDS Homo sapiens (human) SOURCE ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. REFERENCE (bases 1 to 1714) Sasaki, K., Nakatani, Y., Saeki, S., Kawai, H., Nish, T., Nakamura, Y. AUTHORS and Sugano, S. TITLE Novel polypeptide Patent: JP 2001245666-A 1 11-SEP-2001; JOURNAL KYOWA HAKKO KOGYO CO LTD COMMENT Homo sapiens (human) PNJP 2001245666-A/1 PD 11-SEP-2001 06-MAR-2000 JP 2000060548 PF KATSUTOSHI SASAKI, YUKIE NAKATANI, SATOSHI SAEKI, HIROKI KAWAI, PΙ PT TATSUYA NISHI, YUSUKE NAKAMURA, SUMIO SUGANO PΙ C12N15/09, A01H5/00, A01K67/027, A61K39/395, A61K39/395, A61K45/00, PC PC A61P35/00,C07K14/705,C07K16/28,C12N1/15,C12N1/19,C12N1/21, PC C12P21/02, C12Q1/68, C12Q1/68, G01N33/15, G01N33/50, G01N33/53, PC G01N33/566//C12P21/08, (C12P21/02, C12R1:91), C12N15/00, C12N5/00 Location/Qualifiers. **FEATURES** Location/Qualifiers 1..1714 source /organism="Homo sapiens" /mol\_type="genomic DNA" /db xref="taxon:9606" ORIGIN 1 agcacgtaga tcctccctgt catcaggcag agctcttcag tgaggtgggc tcagggaggg 61 ctctgtgcct ccgttcagca gagctgcagc tgctgcccag ctctcaggag gcaagctgga 121 ctccctcact cggctgcagg agcaaggaca gtgaggctca accccgcctg agccatgcca 181 gccaacttca cagagggcag cttcgattcc agtgggaccg ggcagacgct ggattcttcc 241 ccagtggctt gcactgaaac agtgactttt actgaagtgg tggaaggaaa ggaatggggt 301 tecttetaet acteetttaa gaetgageaa ttgataaete tgtgggteet etttgttttt 361 accattgttg gaaactccgt tgtgcttttt tccacatgga ggagaaagaa gaagtcaaga 421 atgacettet ttgtgactea getggeeate acagattett teacaggact ggtcaacate 481 ttgacagata ttaattggcg attcactgga gacttcacgg cacctgacct ggtttgccga 541 gtggtccgct atttgcaggt tgtgctgctc tacgcctcta cctacgtcct ggtgtccctc 601 agcatagaca gataccatgc catcgtctac cccatgaagt tccttcaagg agaaaagcaa 661 gecagggtee teattgtgat egectggage etgtetttte tgtteteeat teecaceetg 721 atcatatttg ggaagaggac actgtccaac ggtgaagtgc agtgctgggc cctgtggcct 781 gacgactect actggacce atacatgace ategtggeet teetggtgta etteateeet 841 ctgacaatca tcagcatcat gtatggcatt gtgatccgaa ctatttggat taaaagcaaa 901 acctacgaaa cagtgatttc caactgctca gatgggaaac tgtgcagcag ctataaccga 961 ggactcatct caaaggcaaa aatcaaggct atcaagtata gcatcatcat cattcttgcc 1021 ttcatctgct gttggagtcc atacttcctg tttgacattt tggacaattt caacctcctt 1081 ccagacaccc aggagcgttt ctatgcctct gtgatcattc agaacctgcc agcattgaat 1141 agtgccatca accccctcat ctactgtgtc ttcagcagct ccatctcttt cccctgcagg 1201 gagcaaagat cacaggattc cagaatgacg ttccgggaga gaactgagag gcatgagatg

1261 cagattctgt ccaagccaga attcatctag accctagggc agtgccagtg ctaggctgag 1321 caccatcage teteceaggt cettgteace tgettgggea egtgeatgga accegageea 1381 acttcacccc accctcgtca ttacctggga gatgcacaag acaaatgttc taatgactgc 1441 atgcactgct taagtattgg ccaacacgaa ctccccagtt attcatgcca gccaggaagg

//

```
1501 aaacgcette etteeceace atteecagee eteetteeca etggecagea eetgaaceea 1561 gtgaacacag geateagtgg teeagggtee tggettggag eeagtgagta gacaggeaag /1621 cagagggac aaaggtaget gggttataca tgaatattet eattacaata ggagaaaata 1681 aaagaettaa ttaageecaa aaaaaaaaaa aaaa
```

```
(360584-42-5) DNA (human G protein-coupled receptor sequence
homolog KAT06734L cDNA plus flanks)
   Score = 3090 Expect = 0.0
   Identities = 1565/1567 (99%)
   Strand = Plus / Plus
            1 gggctcagggagggctctgtgcctccgttcagcagagctgcagctgctgcccagc
   Query:
55
              47 gggctcagggagggctctgtgcctccgttcagcagagctgcagctgctgcccagc
   Subject:
101
           56 tctcaqqaqqcaaqctqqactccctcactcagctgcaggagcaaggacagtgagg
   Query:
110
              102 tctcaggaggcaagctggactccctcactcggctgcaggagcaaggacagtgagg
   Subject:
156
           111 ctcaaccccgcctgagccatgccagccaacttcacagagggcagcttcgattcca
   Query:
165
              157 ctcaaccccgcctgagccatgccagccaacttcacagagggcagcttcgattcca
   Subject:
211
          166 qtqqqaccqqqcaqacqctqqattcttccccaqtqqcttqcactqaaacaqtqac
   Query:
220
              212 gtgggaccgggcagacgctggattcttccccagtggcttgcactgaaacagtgac
   Subject:
266
          221 ttttactqaaqtqqtqqaaqgaaaggaatggggttccttctactactcctttaag
   Query:
275
              267 ttttactgaagtggtggaaggaatggggttccttctactactcctttaag
   Subject:
321
           276 actgagcaattgataactctgtgggtcctctttgtttttaccattgttggaaact
   Query:
330
              322 actgagcaattgataactctgtgggtcctctttgtttttaccattgttggaaact
   Subject:
376
   Query:
           331 ccgttgtgcttttttccacatggaggagaaagaagaagtcaagaatgaccttctt
385
              377 ccgttgtgcttttttccacatggaggagaaagaagtcaagaatgaccttctt
   Subject:
431
           386 tgtgactcagctggccatcacagattctttcacaggactggtcaacatcttgaca
   Query:
              432 tgtgactcagctggccatcacagattctttcacaggactggtcaacatcttgaca
   Subject:
486
           441 gatattaattggcgattcactggagacttcacggcacctgacctggtttgccgag
   Query:
495
```

Expect = 0.0

(450700-57-9) GenBank BD017045

Score = 3090

Length = 1714

Subject:	487 gatattaattggcgattcactggagacttcacggcacctgacctggtttgccgag
Query: 550	496 tggtccgctatttgcaggttgtgctgctctacgcctctacctac
Subject: 596	
Query: 605	551 cctcagcatagacagataccatgccatcgtctaccccatgaagttccttcaagga
Subject:	
Query:	606 gaaaagcaagccagggtcctcattgtgatcgcctggagcctgtcttttctgttct
Subject:	
Query: 715	661 ccattcccaccctgatcatatttgggaagaggacactgtccaacggtgaagtgca
Subject:	
Query:	716 gtgctgggccctgtggcctgacgactcctactggaccccatacatgaccatcgtg
Subject: 816	
Query: 825	771 gccttcctggtgtacttcatccctctgacaatcatcagcatcatgtatggcattg
Subject:	
Query:	826 tgatccgaactatttggattaaaagcaaaacctacgaaacagtgatttccaactg
Subject:	
Query:	881 ctcagatgggaaactgtgcagcagctataaccgaggactcatctcaaaggcaaaa
Subject:	
Query:	936 atcaaggctatcaagtatagcatcatcatcttgccttcatctgctgttgga
Subject:	
Query:	991 gtccatacttcctgtttgacattttggacaatttcaacctccttccagacaccca
1045	

-

Subject: 1037 gtccatacttcctgtttgacattttggacaatttcaacctccttccagacaccca 1091 1046 ggagcgtttctatgcctctgtgatcattcagaacctgccagcattgaatagtgcc Query: 1100 Subject: 1092 ggagcgtttctatgcctctgtgatcattcagaacctgccagcattgaatagtgcc 1146 1101 atcaacccctcatctactgtgtcttcagcagctccatctctttcccctgcaggg Query: 1155 Subject: 1147 atcaacccctcatctactgtgtcttcagcagctccatctctttcccctgcaggg 1201 1156 aqcaaaqatcacaggattccagaatgacgttccgggagagaactgagaggcatga Query: 1210 Subject: 1202 agcaaagatcacaggattccagaatgacgttccgggagagaactgagaggcatga 1211 gatgcagattctgtccaagccagaattcatctagaccctagggcagtgccagtgc Query: 1265 Subject: 1257 gatgcagattctgtccaagccagaattcatctagaccctagggcagtgccagtgc 1311 1266 taggctgagcaccatcagctctcccaggtccttgtcacctgcttgggcacgtgca Query: 1320 Subject: 1312 taggctgagcaccatcagctctcccaggtccttgtcacctgcttgggcacgtgca 1366 1321 tggaacccgagccaacttcaccccaccctcgtcattacctgggagatgcacaaga Query: 1375 Subject: 1367 tggaacccgagccaacttcaccccaccctcgtcattacctgggagatgcacaaga 1421 1376 caaatgttctaatgactgcatgcatgcttaagtattggccaacacgaactcccc Query: 1430 Subject: 1422 caaatgttctaatgactgcatgcatgcttaagtattggccaacacgaactcccc Query: 1431 agttattcatgccagccaggaaggaaacgccttccttccccaccattcccagccc 1485 Subject: 1477 agttattcatgccagccaggaaggaaacgccttccttccccaccattcccagccc 1531 1486 teetteecactqqccaqcacctqaacccagtgaacacaggcattagtggtccagg Query: 1540 Subject: 1532 tccttcccactggccagcacctgaacccagtgaacacaggcatcagtggtccagg 1586 1541 gtcctggcttggagccagtgagtagac 1567 Query: 

Subject: 1587 gtcctggcttggagccagtgagtagac 1613